



PCT/CH 03 / 0075 0

REC'D 2 4 NOV 2003

WIPO

INVESTOR IN PEOPLE

PCT
The Patent Office
Concept House

Cardiff Road

Newport South Wales NP10 8QQ

Rec'd PCT/PTO 11 MAY 2005

I, the undersigned, being an officer duly authorised in accordance with Section 74(1) and (4) of the Deregulation & Contracting Out Act 1994, to sign and issue certificates on behalf of the Comptroller-General, hereby certify that annexed hereto is a true copy of the documents as originally filed in connection with the patent application identified therein.

In accordance with the Patents (Companies Re-registration) Rules 1982, if a company named in this certificate and any accompanying documents has re-registered under the Companies Act 1980 with the same name as that with which it was registered immediately before re-registration save for the substitution as, or inclusion as, the last part of the name of the words "public limited company" or their equivalents in Welsh, references to the name of the company in this certificate and any accompanying documents shall be treated as references to the name with which it is so re-registered.

In accordance with the rules, the words "public limited company" may be replaced by p.l.c., plc, P.L.C. or PLC.

Re-registration under the Companies Act does not constitute a new legal entity but merely subjects the company to certain additional company law rules.

subjects the company to certain additional company law rules.

PRIORITY DOCUMENT

SUBMITTED OR TRANSMITTED IN COMPLIANCE WITH RULE 17.1(a) OR (b)

Signed

Dated

BEST AVAILABLE COPY

n Executive Agency of the Department of Trade and Industry

Patents ADP number (if you know it)

6. If you are declaring priority from one or more earlier patent applications, give the country and the date of filing of the or of each of these earlier applications and (if you know it) the or each application number Country

Priority application number fif you know it)

Date of filing (day/month/year)

 If this application is divided or otherwise derived from an earlier UK application, give the number and the filing date of the earlier application

Number of earlier application

Date of filing (day/month/year)

8. Is a statement of inventorship and of right to grant of a patent required in support of this request? (Answer 'Yes' if:) a) any applicant named in part 3 is not an inventor, or b) there is an inventor who is not named as an applicant, or c) any named applicant is a corporate bady.

See note (d)

yes

Patents Form 1/77

 Enter the number of sheets for any of the following items you are filing with this form.
 Do not count copies of the same document

Continuation sheets of this form

Description 29

Claim(s) 2

Abstract 1

1

Drawing(s)

It is

 If you are also filing any of the following, state how many against each item.

Priority documents

Translations of priority documents

Statement of inventorship and right to grant of a patent (Patents Form 7/77)

Request for preliminary examination and search (Patents Form 9/77)

Request for substantive examination
· (Patents Form 10/77)

Any other documents (please specify)

11.

I/We request the grant of a patent on the basis of this application.

Signature

Date 14/11/02

 Name and daytime telephone number of person to contact in the United Kingdom

Colin Brown (office time) Tel. No: 01/784417721

Warning

After an application for a patent has been filed, the Comptroller of the Patent Office will consider whether publication or communication of the invention should be prohibited or restricted under Section 22 of the Patents Act 1977. You will be informed if it is necessary to prohibit or restrict your invention in this way. Furthermore, if you live in the United Kingdom, Section 23 of the Patents Act 1977 stops you from applying for a patent abroad without first getting written permission from the Patent Office unless an application has been filed at least 6 weeks beforehand in the United Kingdom for a patent for the same invention and either no direction prohibiting publication or communication has been given, or any such direction has been revoked.

Notes

a) If you need help to fill in this form or you have any questions, please contact the Patent Office on 08459 500505

b) Write your answers in capital letters using black ink or you may type them.

o) If there is not enough space for all the relevant details on any part of this form, please continue on a separate sheet of paper and write "see continuation sheet" in the relevant part(s). Any continuation sheet should be attached to this form.

d) If you have answered Yes' Patents Form 7/77 will need to be filed

e) Once you have filled in the form you must remember to sign and date it.

f) For details of the fee and ways to pay please contact the Patent Office





1

Organic Compounds

This invention is concerned with compounds useful for the prevention or suppression of human malodour, in particular human axillary malodour.

It is known that fresh sweat is odourless and that odour is only formed upon contact of sweat with skin bacteria (for example bacteria of the genera of Staphylococcus and Corynebacteria) and it is believed that odourless molecules present in sweat are degraded by bacteria colonising the axilla. It is generally accepted (Labows et. al., Cosmet. Sci Technol. Ser. (1999), 20:59-82) that highly unpleasant malodour is released from fresh sweat mainly by the Corynebacteria genus of bacteria. The principal constituents thought to be responsible for malodour include volatile steroids, volatile sulphur compounds and short-chain, branched fatty acids.

It has been suggested to treat malodour by eradicating the bacteria responsible for causing the odour. Indeed, commercially available cosmetic deodorants often contain antibacterial compounds that generally inhibit the growth of skin microflora.

Antibacterial compounds currently used in deodorant products include, for example Triclosan (2,4,4'-trichloro-2'hydroxy-diphenyl-ether). However, a draw-back to the use of antibacterials is the potential for disturbing the equilibrium of the skin's natural microflora.

Fatty acids, in particular short chain, branched fatty acids are known to play a role in axillary malodour, and are particularly foul smelling components of stale sweat. In copending application PCT/CH0200262 the applicant has disclosed an enzyme that mediates in a process of transforming odourless compounds found in sweat into these malodorous fatty acids. In this co-pending application there is also disclosed a broad class of compounds having activity as inhibitors of the enzyme.

Nevertheless there remains the need to find further compounds displaying good inhibitory properties with respect to the above mentioned enzyme.

2

Accordingly, the invention provided in a first aspect a compound of formula (I)

wherein R is a substituted alkyl, benzyl or allyl residue selected from the group consisting of

nonyl;

4,4,4-trifluoro-propyl;

2-methyl-4-phonyl-butyl;

4-trifluoromethyl-phenyl;

pentafluorophenyl;

4-fluoro-phenyl;

naphthalene-2-yl;

biphenyl-2-yl;

4-tert-butyl-phenyl;

5,5,7,8,8-pentamethyl-5,6,7,8-tetrahydro-naphthalene-2-yl;

5,5,8,8-tetramethyl-5,6,7,8-tetrahydro-naphthalen-2-yl;

1,1,3,3-tetramethyl-indan-5-yl;

styryl;

2,6-dimethyl-hept-1,5-dienyl;

2-(4-tert-Butyl-phenyl)-1-methyl-vinyl;

2-(4-lsopropyl-phenyl)-1-methyl-vinyl;

1-Methyl-3-(2,2,3-trimethyl-cyclopent-3-enyl)-propenyl;

2-(4-Isobutyi-phenyl)-1-methyl-vinyl;

2-(2-isopropyl-phenyl)-1-methyl-ethenyl;

2-phenyl-ethyl;

cyclohexyl-methyl;

2,2-dimethyl-propyl;



3

2-pentafluorophenyl-ethyl;

3-phenyl-propyl; heptyl; and

4-isopropyl-cyclohex-1-enyl.

Compounds of the formula (I) contain chiral atoms and as such they can exist as isomeric mixtures or they may exist as pure stereoisomers. Most preferred are compounds have an S-configuration on the carbon atom in the position alpha to the carboxyl group.

As stated herein above, compounds of the present invention may interact with an enzyme thereby to reduce the enzyme's ability to cleave compounds in sweat leading to release of malodorous acids from odourless fresh sweat. That enzyme, described in the afore-mentioned co-pending application, was isolated from the bacteria of the genus Corynebacteria that can be found colonising the axilla, in particular certain Corynebacteria sp., more particularly Corynebacteria striatum Ax 20 which has been submitted on the 26, April 2001 to the International Depository Authority DSMZ-Deutsche Sammlung von Mikrooganismen und Zellkulturen GmbH, D-38124 Braunschweig. The Accession Number provided by the International Depository Authority is DSM 14267. The enzyme was found to occur intracellularly and can be released from the cells by mechanical disruption of the cell envelope. Thus, it may be isolated from cellular extracts obtained from wild-type bacterial strains, especially from strains of Corynebacteria isolated from the human axilla, in particular Corynebacterium striatum Ax 20. In the alternative, it may be produced by recombinant means which are well known to persons skilled in the art.

The amino acid sequence of this enzyme is set forth in SEQ ID No. 1 and a nucleic acid sequence encoding for this enzyme is set forth in SEQ ID No. 2, both of which sequences are shown below.

Compounds of the present invention display inhibition of the enzyme at concentrations of about 1 to 500,nanomolar more particularly from nanomolar to tens of nanomolar concentration in vitro, e.g. from 10 to 20 nanomolar. Furthermore, having regard to the

4

lipophilicity of the residue R, the compounds are adapted to penetrate the cell walls of the enzyme-producing bacteria, as such, they are efficaceous in vivo.

Indeed, the nature of the residue R appears to influence the ability of compounds to penetrate the cellular walls of different bacteria colonising the axilla and which are implicated in malodom production. For example, other strains of Corynebacteria, or bacteria of the genus Staphylococci found in the microflora of the axilla also produce related enzymes that themselves mediate in biochemical reactions wherein L-glutamine derivatives are cleaved at N_{α} . The compounds of the present invention may interfere in collular processes of a wide variety of bacterial strains thereby resulting in the suppression or prevention of malodour from these sources.

The in vitro activity of the compounds as inhibitors may be measured in terms of either their IC_{50} values or their Ki values, both of which measures are well known to the person skilled in the art. As is well known, the IC_{50} value provides the concentration of an inhibitor needed to reduce enzyme velocity by half at a given substrate concentration. This value is dependent on the affinity of the substrate for the enzyme which is reflected in the value K_m of the substrate. In this way, the Ki value may be determined for a given substrate and a given substrate concentration by measuring IC_{50} and then calculating according to the following formula

$$K_{l} = \frac{lC_{50}}{1 + \frac{[Substrate]}{K_{m}}}$$

The uptake of the compounds in bacterial cells and the inhibition of the enzyme contained therein may be measured using an assay based on stationary-phase living cells. Thus, cells may be incubated along with inhibitory compound or compounds, and the substrate (i.e. the material found in sweat, which when cleaved by the enzyme forms the malodorous acids), and the release of acids may be measured at various inhibitor concentrations. By comparing IC_{50} values obtained with the living cells with the IC_{50}

5

values obtained with the isolated enzyme the ease of penetration of the compounds into the bacterial cells can be assessed.

Compounds of the present invention may be added to any cosmetic and personal care products such as sticks, roll-ons, pump-sprays, aerosols, deoderant soaps, powders, solutions, gels, creams, sticks, balms and lotions to enhance the deoderising effect of these products. Preferably, a compound of the present invention may be employed in said products in amounts of about 0.01 to 0.5% by weight.

The above-mentioned products, in addition to the inhibitors, may comprise anti-bacterial agents known in the art, e.g. Triclosan. The products may also comprise dermatologically acceptable ingredients such as are commonly used in these types of product. Examples of such additional ingredients include fragrances, colorants, opacifiers, buffers, antioxidants, vitamins, emulsifiers, UV absorbers, silicones and the like. As is also well known, all products can be buffered to the desired pH.

In addition to the inhibitor, a deodorant cologne may comprise ethanol and fragrance. Fragrance may be present from 1 to 10% and the ethanol can be present to make up the mass to 100%.

Additional ingredients in a typical ethanol-free decodorant stick may include polyols, such as propylene glycol; derivatives thereof, such as propylene-glycol-3-myristyl ether (Witconol APM); water; a surfactant such as sodium stearate; and a fragrance. The polyol may be present in an amount of 30 to 40%; the derivatives of the polyol likewise may be present at about 30 to 40%; water may be present to about 10 to 20%; the surfactant may be present to 5 to 10%; and the fragrance may be present in an amount mentioned above.

A typical antiperspirant stick might contain as additional ingredients such as Ethylene Glycol Monostearate (e.g. from 5 to 10%); Shea butter (e.g. from 3 to 5%); Neobee 1053 (PVO International) (e.g. from about 12 to 15%); Generol 122 (Henkel) (e.g. from

6

824 29 26:

about 3 to 7%); Dimethicone (DC 345)(e.g. from 30 to 40%); aluminium sesquichlorohydrate (e.g. from about 15 to 20%); and a fragrance, e.g. from 1 to 10%.

An antiperspirant aerosol may contain ethanol, e.g. from about 10 to 15%; zirconium aluminium tetrachlorohydrate, e.g. from about 3 to 5%; Bentone 38, e.g. from about 1 to 2%; fragrance in an amount aforementioned; and a hydrocarbon propellant, e.g. S-31 up to 100%.

An antiperspirant pump composition may contain aluminium sesquichlorohydrate, e.g. from 15 to 25%; water, e.g. from 50 to 60%; Triton X-102 (Union carbide), e.g. from 1 to 3%; dimethyl Isosorbide (ICI), e.g. from 15 to 25 %; and a fragrance in an amount as aforementioned.

All percentages mentioned above are in wt %.

Accordingly, the present invention relates to the use of the compounds and/or compositions containing same for the elimination or suppression of malodour. The invention also relates to compositions comprising an odour suppressing quantity of an inhibitor of the enzyme and dermatologically acceptable vehicles which are generally well known in the art of cosmetic and personal care products.

The invention also provides in another of its aspects, a method of suppressing axillary malodour comprising the step of providing a composition for application to a person in need of treatment, said composition containing an inhibitor compound and dermatologically acceptable vehicle therefor, said compound being selected from one or more compounds described above.

A compound of formula (I) may be prepared according to synthetic protocols as set out in detail below with reference to Scheme 1, Scheme 2 and the Examples

7

- A) 5 equivalents (eq.) HP(OTMS)2, for 2 h at 130°C.
- B) 1 eq alkylphosphinic acid 3, 5 eq HMDS for 3 h at 130°C, then 1 eq acrylate 1 for 4 h at 130°C, then EtOH at 70°C. The product is obtained in quantitative yield.
- C) 1 eq benzylic or allylic bromide (6 or 8), 3 eq BSA, at 25°C. The product is obtained in quantitative yield.
- D) 1 N LiOH / EtOH 1:1 (weight-equivalents) for I day at 25°C to give the product in quantitative yield.
- E) 2.2 eq (iPr)₃SiH in TFA at 25°C for 3 h to provide compounds of the present invention.

The acrylate starting material (1) may be formed according to a method described in copending application PCT/CH0200262 and is set forth in Scheme 3 below.

The alkyl, benzyl or allyl halides are either commercially available or may be formed from commonly available starting materials according to synthetic protocols known per se and set out in Scheme 2 below.

8

- F) I eq Br2 at 170°C for 4 h.:
- G) 2 eq Pyridine, 1.2 eq PPh3; 1.2 eq Iodine at 0°C for 2 h.
- H) 0.35 eq NaBH4, MeOH for 2 h at 0°C, to provide the allyl alcohols in quantitative yield.
- I) Et₂O, 0.4 eq PBr₃ for 5 h at 0°C to provide allyl bromides in quantitative yields.
- K) 3-5 eq HP(OTMS)2 in CH2Cl2, for 16 h.
- L) 2 eq NaH₂PO₂(H₂O), 1 eq BEt₃, MeOH for 6 h at 25°C.

9

There now follows a series of Examples that serve to illustrate the invention.

Synthesis Examples 1 to 9

The following compounds are formed according to the following syntheses:

- 5a 4-Carbamoyl-2-(decyl-hydroxy-phosphinoylmethyl)-butyric acid
- 5b 4-Carbamoyl-2-(hydroxy-(4,4,4-trifluoro-butyl)-phosphinoylmethyl]-butyric acid
- 5c 4-Carbamoyl-2-[hydroxy-(3-methyl-5-phenyl-pentyl)-phesphinoylmethyl]-butyric acid
- 5d 4-Carbamoyl-2-[hydroxy-(4-trifluoromethyl-benzyl)-phosphinoylmethyl]-butyric acid
- 5e 4-Carbamoyl-2-(hydroxy-pentafluorophenylmethyl-phosphinoylmethyl)-butyric acid
- 5f 4-Carbamoyl-2-[(4-fluoro-benzyl)-hydroxy-phosphinoylmethyl]-butyric acid
- 5g 4-Carbamoyl-2-(hydroxy-naphthalen-2-ylmethyl-phosphinoylmethyl)-butyric acid
- 5h 2-(Biphenyl-2-ylmethyl-hydroxy-phosphinoylmethyl)-4-carbamoyl-butyric acid
- 5i 2-[(4-tert-Butyl-benzyl)-hydroxy-phosphinoylmethyl]-4-carbamoyl-butyric acid
- 5j 4-Carbamoyl-2-[hydroxy-(5,5,7,8,8-pentamethyl-5,6,7,8-tetrahydro-naphthalen-2-ylmethyl)-phosphinoylmethyl]-butyric acid
- 5k 4-Carbamoyl-2-[hydroxy-(5,5,8,8-tetramethyl-5,6,7,8-tetrahydro-naphthalen-2-ylmethyl)-phosphinoylmethyl]-butyric acid
- 51 4-Carbamoyl-2-[hydroxy-(1,1,3,3-tetramethyl-indan-5-ylmethyl)-phosphinoylmethyl]-butyric acid
- 5m E-4-Carbamoyl-2-[hydroxy-(3-phenyl-allyl)-phosphinoylmethyl]-butyric acid
- 5n E-4-Carbamoyl-2-[(3,7-dimethyl-octa-2,6-dienyl)-hydroxy-phosphinoylmethyl]-butyric acid

1

10

- 50 E-2-{[3-(4-tert-Butyl-phenyl)-2-methyl-allyl]-hydroxy-phosphinoylmethyl}-4-carbamoyl-butytic acid
- 5p E-4-Carbamoyl-2-{hydroxy-[3-(4-isopropyl-phenyl)-2-methyl-allyl]-phosphinoylmethyl}-butyric acid
- 5q E-4-Carbamoyl-2-{hydroxy-[2-methyl-4-(2,2,3-trimethyl-cyclopent-3-enyl)-but-2-enyl]-phosphinoylmethyl}-butyric acid
- 5r E-4-Carbamoyl-2-{hydroxy-[3-(4-isobutyl-phenyl)-2-methyl-allyl]-phosphinoylmethyl}-butyric acid
- 5s E-4-Carbamoyl-2-{hydroxy-[3-(2-isopropyl-phenyl)-2-methyl-allyl]-phosphinoylmethyl}-butyric acid
- 5t 4-Carbamoyl-2-[hydroxy-(3-phenyl-propyl)-phosphinoylmethyl]-butyric acid
- 5u 4-Carbamoyl-2-[(2-cyclohexyl-ethyl)-hydroxy-phosphinoylmethyl]-butyric acid
- 5v 4-Carbamoyl-2-[(3,3-dimethyl-butyl)-hydroxy-phosphinoylmethyl]-butyric acid
- 5w 4-Carbamoyl-2-[hydroxy-(3-pentafluorophenyl-propyl)-phosphinoylmethyl]butyric acid
- 5x 4-Carbamoyl-2-[hydroxy-(4-phenyl-butyl)-phosphinoylmethyl]-butyric acid
- 5y 4-Carbamoyl-2-(hydroxy-octyl-phosphinoylmethyl)-butyric acid
- 5z 4-Carbamoyl-2-[hydroxy-(4-isopropyl-cyclohex-1-enylmethyl)phosphinoylmethyl]-butyric acid

Structures of these compounds are set out below:

11

12

The following Examples are described with reference to Scheme 1 and Scheme 2.

Example 1

Preparation of 2-Hydroxyphosphinoylmethyl-4-(trityl-carbamoyl)-butyric acid ethyl ester 2: (Step A of Scheme 1)

The synthesis is carried out analogously to L.A.Reiter, B.P.Jones, J.Org.Chem. 62, 2808 (1997) or W.P.Malachowski, J.K.Coward, J.Org.Chem. 59, 7625 (1994).

In a 500mL flask equipped with a septum and a condenser, 25 g (0.3 mol) ammonium phosphinate and 49 g (0.3 mol) HMDS are heated under N₂ at 110°C for 3.5 h. The reaction mixture is cooled to 5°C where 25 g acrylate 1 (the synthesis of which is set out

13

in Scheme 3 above) in 150 ml dichloromethane are added. The mixture is stirred for 16 h at room temperature. Work-up: 1 N HCl and CH₂Cl₂ are added. The organic phase is washed with 1 N HCl, the combined acidic phases are re-extracted with CH₂Cl₂. The combined organic phases are dried over MgSO₄, evaporated under reduced pressure and dried at 50°C under high vacuum yielding 28.8 g (99.9%) of phosphinic acid 2.

Yield: 99.9%

M.p.: 152-154°C (white solid).

Purity: 89 % (31P-NMR)

³¹P-NMR (CDCl₃, 400MHz): 32.0 ppm (s).

MS (ESI neg.): 957 [2M - H], 478 [M - H].

¹H-NMR (CDCl₃, 400MHz): 1.2 (t, 3H), 1.8 (m, 1H), 1.9 (2H), 2.05 (m, 1H), 2.25 (m, 1H), 2.7 (m, 1H), 4.1 (q, 2H), 6.3 - 7.7 (d, 1H, P-H, J = 560 Hz), 6.9 (s, 1H, NH), 7.2 (15H, trityl-H), 8.2 (1H, P-OH).

¹³C-NMR (CDCl₃, 400MHz): 14.2 (CH₃), 28.7 (d, CH₂), 30.8, 31.8 (d, P-CH₂), 34.3 (s, CH₂), 38.2 (CH), 61.2 (OCH₂), 70.5 (Ph₃C), 127.0 (3C, Trityl-CH), 127.9 (6C, Trityl-CH), 128.7 (6C, Trityl-CH), 144.6 (3C, Trityl-C), 171.0 (C=O), 174.0 (C=O).

Example 2

Preparation of 4-Carbamoyl-2-[hydroxy-(4,4,4-trifluoro-butyl)-phosphinoylmethyl]-butyric acid 4b: (Step B of Scheme 1)

The synthesis is based on Boyd, E.A.; Regan, A.C.; Tetrahedron Lett 24, 4223 (1994) as described in the previously filed patent 30012/PCT.

0.3 g (1.7 mmol) (4,4,4-trifluoro-butyl)-phosphinic acid 3b (0.3 g, 1.7 mmol) are dissolved in HMDS (1.4 g, 8.5 mmol) at room temperature and heated at 130°C for 4 h. At 80°C acrylate 1 (0.7g, 1.7 mmol) is added and the reaction mixture heated at 130°C for 16 h. Ethanol is added at 60°C and the mixture refluxed for 30 min. The solvents are removed under reduced pressure and the residue is dried at 50°C under high vacuum for 8 h to yield 0.9 g (89%)of 4b.

14

. Yield: 89 %.

Purity: ~80 % ('H-NMR)

³¹P-NMR (CDCl₃, 400MHz): 45.3 ppm (s).

MS (ESI neg.): 670 (8% [M + NaOAc - H), 588 (100% [M - H]).

¹H-NMR (CDCl₃, 400MHz): 1.2 (t, 3H), 1.55 (m, 2H), 1.7 (m, 1H), 1.8 (3H), 1.95 (m, 1H), 2.15 (3H), 2.2 (m, 1H), 2.4 (m, 1H), 2.75 (m, 1H), 4.1 (q, 2H), 6.8 (s, 1H, NH), 7.2 (Trityl-H).

Whereas this synthesis is described with reference to the "R" residue relating to compound 5b (above), this synthesis is carried out for the preparation of other alkyl or aralkyl phosphinic acids whose "R" residues correspond to the compounds 5a, 5c, 5f 5i, 5n, 5t, 5u, 5v, 5w, 5x, 5y and 5z (all described above).

Example 3:

Preparation of 2-[Hydroxy-(1,1,3,3-tetramethyl-indan-5-ylmethyl)-phosphinoylmethyl]-4-(trityl-carbamoyl)-butyric acid ethyl ester (41) (Step C of Scheme 1)

This synthesis is based on a similar BSA-mediated alkylation of an alkylphosphinic acid described by L.A.Reiter, B.P.Jones, *J.Org.Chem.*62, 2808 (1997).

In a 100mL flask equipped with a septum and a condenser, monoalkylphosphinic acid 2 (3g, 6.4 mmol) is dissolved in dry CH₂Cl₂ (20 ml). 5-Bromomethyl-1,1,3,3-tetramethyl-indan 6I (1.9g, 7 mmol) and BSA (3.9g, 19 mmol) are added and the mixture is stirred 72 h at 25°C. Work-up: The mixture is poured on 1N HCl. The organic phase is washed with 1N HCl, the combined acidic phases are re-extracted with 1N HCl. The combined organic phases are dried over MgSO₄, evaporated under reduced pressure and dried at 50°C under high vacuum to give 4.77 g (100%) of the bisalkylated phosphinic acid 41.

Yield: 100%

Purity: 82 % (2) P-NMR) -

³¹P-NMR (CDCl₃, 400MHz): 53,8 ppm (s).

15

MS (ESI neg.): 1329 (10% [2M – H]), 664 (100% [M – H]), 494 (30%). 1 H-NMR (CDCl₃, 400MHz): 1.2 (t, 3H), 1.3 (14H), 1.7 (m, 1H), 1.9 (d, 2H, P-CH₂), 2.1 (m, 1H), 2.25 (2H), 2.7 (m, 1H), 3.0 (d, 2H, P-CH₂), 4.1 (q, 2H, OCH₂), 6.85 (s, 1H, NH), 7.2 (15H, trityl-H), 8.4 (s, 1H, P-OH).

Whereas this synthesis is described with reference to the "R" residue relating to compound 51 (above), this synthesis is carried out for the preparation of other benzylic or allylic phosphinic acids whose "R" residues correspond to the compounds 5d, 5e, 5g, 5 h, 5j, 5k, 5m, 5o, 5p, 5q, 5r and 5s (all described above).

Example 4

Hydrolysis / detritylation of bisalkyl-phosphinoyl compounds 4 to the P-alkylated 2-(Alkyl-hydroxy-phosphinoylmethyl)-4-carbamoyl-butyric acids 5: (Steps D and E in Scheme 1)

A compound 4 (1g, 1.9 mmol) is solubilized in 9mL ethanol. 9mL 1N LiOH (5eq) are added. The reaction mixture is stirred 3h at room temperature. 1N HCl is added until pH=1 and BtOH is removed under reduced pressure. The aquous residue is extracted with CH₂Cl₂. The organic layers are dried over MgSO₄. Evaporation of the combined organic layers gives 0.75 g of hydrolyzed intermediate. Complete Hydrolysis is checked by ¹H-NMR, the structure can be further confirmed by ³¹P-NMR and MS (BSI neg.). The material is subjected to detritylation without further purification:

The above acid (0.5g, 1mmol) is solubilized in 10mL TFA. 0.35 g (2.2mmol) iPr3SiH are added via syringe. The reaction mixture is stirred for 2 h at room temperature. TFA is removed under reduced pressure and the reaction mixture co-evaporated with toluene. The crude product is dissolved in 7 ml water at 50°C and filtered over a RPC-18 syringe. The filtrate is co-evaporated with Toluene giving 0.1 g of 5 (42%).

All of the compounds 5 described above are obtained with purities: 85-90% (31P-NMR).

Example 5

16

Preparation of 7-Bromomethyl-1,1,2,4,4-pentamethyl-1,2,3,4-tetrahydro-naphthalene 6j: (Step F of Scheme 2)

The solventless bromination of substituted toluenes at higher temperatures is based on well known literature methods, e.g. I.B.Shoesmith, A.Mackie, *J.Chem.Soc.*, 300 (1936) or J.B.Baker, *J.Chem.Soc.*, 1448 (1936).

In a 100mL flask equipped with thermometer, septum and a condenser 21.6 g (0.1 mol) 1,1,2,4,4,7-hexamethyl-1,2,3,4-tetrahydro-naphthalene (prepared as described by Wood, T. F.; Easter, W. M., Jr.; Carpenter, M. S.; Angiolini, J. Org. Chem.28, 2248 (1963)) are heated to 170°C. 16 g (0.1 mol) Bromine are added and the reaction mixture is stirred for 5 h at 170°C. The content of the flask is fractionated over a Vigreux column (110°C, 0.04 Torr) giving 17 g (58%) 6j as a colorless liquid.

Yield: 58%.

GC-purity: 71%

Impurities:

7% 1,1,2,4,4,7-hexamethyl-1,2,3,4-tetrahydro-naphthalene

14% 7-Dibromomethyl-1,1,2,4,4-pentamethyl-1,2,3,4-tetrahydro-naphthalene

GC/MS: 294 / 296 (5%, [M][†]), 279 / 281 (10%, [M – CH₃][†]), 215 (100%, [M – Br][†], 201 (65%, [M – CH₂Br][†]), 157 (55%).

¹H-NMR (CDCl₃, 400MHz): 0.95 (d, 3H), 1.05 (s, 3H), 1.25 (s, 3H), 1.3 (s, 3H), 1.35 (s, 3H), 1.4 (dd, 1H, A), 1.65 (dd, 1H, B), 1.85 (m, 1H, CH), 4.5 (s, 2H, CH₂Br), 7.15 (d, 1H, Ar-H), 7.25 (d, 1H, Ar-H), 7.35 (s, 1H, Ar-H).

¹³C-NMR (CDCl₃, 400MHz): 16.8 (CH₃), 25.0 (CH₃), 28.5 (CH₃), 31.9 (CH₃), 32.3 (CH₃), 34.3 (CH₂), 34.4 (C), 34.5 (CH), 37.8 (C), 43.5 (CH₂Br), 126.2, 127.0, 127.7 (Ar-CH), 134.7, 145.2, 146.5 (Ar-C).

Example 6

Preparation of (5-Iodo-3-methyl-pentyl)-benzene 7c: (Step G of Scheme 2)

17

The synthesis proceeded according to the well known literature method, see for example M.Ishizaki, O.Hoshino et al., *Tetrahedron* 57, 2729 (2001).

5 g (28 mmol) Phenoxanol (3-Methyl-5-phenyl-pentan-1-ol) are dissolved in 200 ml dichloromethane under nitrogen and stirring. Triphenylphosphine (8.8 g, 34 mmol) and 4.2 g (53 mmol) pyridine are added at 25°C. At 0°C iodine (8.5 g, 34 mmol) are added. After 2 h stirring at 0°C the reaction mixture is poured on ice-cooled 1 N HCl and extracted with dichloromethane. The combined organic phases are washed with 10% Na₂S₂O₃, sat. NaHCO₃ and sat. NaCl. Drying over MgSO₄ and evaporation gives 17g of a residue which is triturated with hexane and filtered over a 5 cm Silicagel pad. The filtrate is evaporated under reduced pressure and dried under high vacuum giving 7.1 g (88%) of 7c as an colorless oil.

Yield: 88%.

Purity: > 95% (GC, NMR)

GC/MS: 288 (5%, $[M]^{\dagger}$), 161 (10%, $[M-I]^{\dagger}$),

119 (5%), 105 (20%, [PhCH₂CH₂]⁺), 91(100%, [[PhCH₂]⁺).

¹H-NMR (CDCl₃, 400MHz): 0.95 (d, 3H, CH₃), 1.45 (m, 1H), 1.6 (3H), 1.9 (m, 1H),

2.6 (2H, PhCH₂), 3.2 (2H, CH₂I), 7.2 (5H, A₁H).

¹³C-NMR (CDCl₃, 400MHz): 4.9 (CH₂), 18.7 (CH₃), 33.3 (CH₂), 33.7 (CH₃), 38.2 (CH₂), 40.9 (CH₂), 125.8, 128.3, 128.4 (ArCH), 142.5 (ArC).

Example 7

Preparation of E-1-(3-Bromo-2-methyl-propenyl)-4-tert-butyl-benzene 80: (Steps H and I of Scheme 2)

Both steps are well known literature methods, e.g. R.M.Coates, D.A.Ley, P.L.Cavender, J.Org.Chem.43, 4918 (1978).

20 g (0.1 mol) E-3-(4-tert-Butyl-phenyl)-2-methyl-propenal (prepared as described at Y.Crameri, P.A.Ochsner, P. Schudel, US 4435585, Givaudan (1982)) are added to a

18

stirred solution of 1.2 g (32 mmol) of sodium borohydride in 20 ml of methanol at 0°C. After 2 h at room temperature quantitative conversion is checked by TLC. The reaction mixture is poured onto 40 ml saturated sodium chloride and extracted with tert-butyl methyl ether. Drying over magnesium sulfate and evaporation of the solvent gives the crude allylic alcohol (19.2 g, 94%), which is transferred to the following bromination without further purification.

5 g (24 mmol) of the crude allylic alcohol are dissolved in 35 ml of dry diethyl ether under nitrogen. At 0°C phosphorus tribromide (0.95 ml, 10 mmol) is added via syringe. The reaction is stirred at 0°C for 4 h, poured onto ice and extracted three times with diethyl ether. The organic layer is washed with saturated NaHCO₃ and dried over MgSO₄. The solvent is removed under reduced pressure giving the crude allylic bromide 5.7 g (87%) of 80.

Yield: 5.7 g (82% from the aldehyde).

Purity: > 95% (GC, NMR)

GC/MS: 266 / 268 (3%, [M]^T), 251 (1%, [M – CH₃][†]), 188 (25%, [M – Br][†]), 173 (55%, [M – Br – CH₃][†]), 157 (10%), 131 (55%), 115 (22%); 91 (16%), 57 (100%).

¹H-NMR (CDCl₃, 400MHz): 1.3 (s, 9H, tBu-CH₃), 2.0 (s, 3H, CH₃), 4.15 (s, 2H, CH₂Br), 6.6 (s, 1H, =CH), 7.22 (d, 2H, Ar-H), 7.35 (s, 2H, Ar-H).

¹³C-NMR (CDCl₃, 400MHz): 16.6 (CH₃), 31.3 (3C, tBu-CH₃), 34.6 (tBu-C), 42.6 (CH₂), 125.2, 128.7 (ArCH), 130.0 (=CH), 133.7, 134.0 (Ar-C), 150.2 (=C)

Example 8

Preparation of (3,7-Dimethyl-octa-2,6-dienyl)-phosphinic acid 3n: (Step K of Scheme: 2)

Synthesis is based on the publication of E.A.Boyd, A.C.Regan, *Tetrahedron Letters*, 24, 4223 (1994). Recently it has been been found that a 3-5 eq excess of the reagents HMDS and ammonium phosphinate improves the yield of alkyl phosphinicacid 3 formation when alkyl iodides 7 are employed (H.An et al., *J.Org.Chem.66*, 2789 (2001), S.Chem, J.K.Coward, *J.Org.Chem.63*, 502 (1998)) and that an excess of HMDS



19

and NH₄⁺PO₂H₂⁻ prevents also bis-alkylation using benzylic- or allylic bromides 6 or 8 (S.Depréle, J.-L.Montchamp, *J.Org. Chem.* 66, 6745 (2001)).

In a 750mL flask equipped with septum, thermometer and condenser, ammonium phosphinate (25 g, 0.3 mol) and HMDS (51 g, 0.32 mmol) are heated under N₂ at 110°C for 3 h. The reaction mixture is cooled to 0°C. 300 mL dried CH₂Cl₂ are added followed by the addition of geranyl bromide (13.1 g, 60 mmol). The mixture is stirred for 16 h at room temperature. 12 g methanol are added and the fine suspension is filtered over a double filter layer. The filtrate is concentrated under reduced pressure. 10% Na₂CO₃ and text-butyl methyl ether are added, the phases are separated and the alkaline layer purified with text-butyl methyl ether. The alkaline layer is treated with conc. HCl until pH = 1 and is 3 times extracted with dichloromethane. Drying of the dichloromethane layer over MgSO₄ and evaporation give 13.7 g (81%) of 3n as an orange oil.

Yield: 13.7 g (81%),

Purity: 77% (31P-NMR)

³¹P-NMR (CDCl₃, 400MHz): 34.9 ppm (s).

MS (ESI neg.): $403 (10\% [2M - H]^{+})$, $201 (100\%, [M-H]^{+})$.

¹H-NMR (CDCl₃, 400MHz): 1.2 (d, 1H), 1.6 (s, 3H, CH₃), 1.6 (6H, 2 CH₃), 2.1 (4H, CH₂CH₂), 2.6 (dd, 2H, P-CH₂), 5.1 (1H, =CH), 5.15 (1H, =CH), 6.22 and 7.6 (d, 1H, J = 548 Hz, P-H), 11.8 (s, 1H, POH).

¹³C-NMR (CDCl₃, 400MHz): 16.5 (CH₃), 17.6 (CH₃), 25.6 (CH₃), 26.4 (CH₂), 30.4 and 29.5 (d, P-CH₂), 39.7 (CH₂), 110.5 (=CH), 123.7 (=CH), 131.7 (=C), 141.9 (=C).

Example 9

Preparation of (3-phenyl-propyl)-phosphinic acid 3t: (Step L of Scheme 2)

The method has been described by S. S.Deprele, J.-L.Montchamp, J.Org. Chem. 66, 6745 (2001).

To a solution of NaH₂PO₂(H₂O) (13.2 g, 0.125 mol) and allylbenzene (6.6 g, 56 mmol) in methanol (250 ml) is added triethylborane (1M in THF, 50 ml, 50 mmol) at room



20

temperature in an open 500 ml flask. The solution is stirred for 2 h at room temperature. The reaction mixture is concentrated under reduced pressure. 100 ml saturated KHSO₄ are added to the residue followed by extraction (200, 100 and 70 ml) with ethyl acetate. To the combined ethyl acetate phases are added 40 ml 10% Na_2CO_3 . Under vigorous stirring and dropwise addition of conc. NaOH the biphasic mixture is adjusted to pH = 10. The organic phase is separated and the alkaline phase adjusted to pH = 2 by addition of conc. HCl. Extraction with chloroform (3 x 100 ml), drying of the combined organic layer over MgSO₄ and evaporation gives 5.4 g of crude 3t (61%).

Yield: 61%

Purity: 84% (31 P-NMR)

³¹P-NMR (CDCl₃, 400MHz): 38.2 ppm (s).

MS (ESI neg.): $265 (6\% [M-H-NaOAc]^{+})$, 183 (100%, $[M-H]^{+}$).

¹H-NMR (CDCl₃, 400MHz): 1.75 (m, 2H, CH₂), 1.9 (m, 2H, CH₂), 2.76 (t, 2H, PhCH₂), 6.4 and 7.7 (d, 1H, J = 548 Hz, P-H), 11.4 (s, 1H, POH).

Application Examples

Example 1,

Isolation of new malodour acid and precursors thereof from human sweat:

Fresh axilla secretions are sampled from human panellists by washing the axilla with 10% ethanol. The samples are extracted with MTBE to remove interfering lipids. The hydrophilic phases obtained from the washings from several individuals are then pooled. This material is practically odourless, but upon hydrolysis of sub-samples with 1 M NaOfl, it produced typical axilla malodour. To identify the malodour volatiles, hydrolysed sub-samples are extracted and concentrated by solid phase extraction and then analysed by GC-sniff. Peaks that are rated as having a strong odour and closely related to axilla malodour are analysed by GC-MS. The samples contained one particular peak of an acid very typical of axilla malodour. Based on the MS data the most probable structure of this peak is 3-hydroxy-3-methyl-hexanoic acid. This assumption is verified by synthesising this latter compound and comparing its spectra

14

";

Von: Givaudan / Fragrance Research

30076GB

21 .

and retention times to the GC-MS data of the major malodour peak in the GC-sniff analysis. This malodour compound is structurally related to the known sweat malodour acid 3-methyl-2-hexenoic acid, and it is transformed into this latter compound by dehydration upon prolonged incubation.

To identify the precursor for this acid, the pooled non-hydrolysed sample is separated on a Superdex gel filtration column (Pharmacia, Uppsala, Sweden) using NH₄CO₃/NaCl as the elution buffer. Individual fractions of this separation step are tested for the content of a malodour precursor by hydrolysis with 1 M NaOH. One fraction developed strong malodour upon hydrolysis and this malodour could be attributed to the release of 3-hydroxy-3-methyl-hexanoic acid by GC-MS analysis. This fraction is subjected to LC-MS analysis. It contained one major mass peak of 274 Da and an additional peak at 256 Da. The mass spectrum of the former peak suggested a compound where the 3-hydroxy-3-methyl-hexanoic acid is linked to one molecule of L-glutamine (i.e. Nα-3-hydroxy-3-methyl-hexanoyl-L-glutamine), while the second peak could, based on its mass, correspond to the dehydrated analogue Nα-3-methyl-2-hexenoyl-L-glutamine. Nα-3-hydroxy-3-methyl-hexanoyl-L-glutamine is then synthesised and its MS spectrum and retention time in the LC-MS-analysis compared to and found identical with the compound isolated from natural sweat.

Example 2

Isolation of axilla bacteria having the ability to cleave the malodour precursor compound

The axillary flora of 8 panellists is isolated with the detergent-scrub method: A 6 cm² area of the axilla is scrubbed with a phosphate buffer at pH 7 containing 1% Tween 80. The samples are spread-plated on tryptic soy agar amended with 5g/L of Tween 80 and 1 g/L of lecithin. Single isolates obtained after 48 h incubation are subcultured and characterised. A total of 24 individual strains are identified based on colony and cell morphology, gram-reaction, lipophilic growth, lipase reaction and API identification kits (bioMerieux, France; coryneforms with the API coryne kit and cocci with the ID

i. .

22

Staph 32 kit). The strains are grown overnight in a liquid medium (Mueller-Hinton amended with 0.01% Tween 80), harvested by centrifugation and resuspended to a final OD_{600} of 1 in a semi-synthetic medium (Per litre; 3 g KH₂PO₄, 1.9 g K₂HPO₄, 0.2 g yeast extract, 0.2 g MgSO₄, 1.4 g NaCl, 1 g NH₄Cl, 10 mg MnCl₂, 1 mg Fe₃Cl₂, 1 mg CaCl2). Aliquots of this stationary culture are then amended with a final concentration of 500ppm of N_{α} -3-hydroxy-3-methyl-hexanoyl-L-glutamine (5% stock solution dissolved in methanol). After 24 h incubation (with shaking at 300rpm; 36°C) the -samples are extracted and the amount of released 3-hydroxy-3-methyl-hexanoic acid is :-determined by capillary GC. Table 1 gives the results for a subset of the strains tested. From these results it appears that among the Corynebacteria isolated from the axilla some, but not all, are able to release 3-hydroxy-3-methyl-hexanoic acid from the synthetic precursor. The Corynebacteria which are able to conduct this biochemical reaction may be found in the group of the lipophilic and in the group of the nonlipophilic Corynebacteria. Therefore, a specific enzyme only present in some bacterial strains seems to be responsible for this cleavage. Since it releases axilla malodour the putative enzyme is named AMRB, which stands for 'axillary malodour releasing enzyme'. Apparently the tested Staphylococci are not able to catalyse this reaction, which is in agreement with the observation; that only subjects with an axilla flora dominated by Corynebacteria produce the most typical axilla malodour (Labows et. at.; Cosmet. Sci Technol. Ser. 1999, 20:59-82). However, when $N\alpha$ -lauroyl-L-glutamine is used as substrate in the same experiment, it is found that also other Corynebacteria and some Staphylococci can release lauric acid from this substrate. It therefore appears, that most axilla bacteria have a related enzyme, but that many can only release straight fatty acids which make a minor contribution to typical axilla malodour.

Table 1. Cleavage of the natural malodour precursor by axilla bacteria.

Isolate	Species assignment	Lipophilic	3-hydroxy-3-methyl-hexanoic acid released (ppm)			
Axl	Staphylococcus capitis	-	O (Phitt)			
Ax6	Staphylococcus epidermidis Micrococcus luteus		0			
		•	0			
WYD	Corynebacterium bovis	+	0			



Ax7	Corynebacterium group G		+	٠.		0
Ax15	Corynebacterium jeikelum		+			37.4
Ax19	Corynebacterium jelkeium	•	+		•	105.1
Ax20	Corynebacterium striatum		-			262.7

(*) Corynebacteria isolated from the human axilla may be separated into two classes based on their requirement for a source of fatty acids in the growth medium.

Example 3

Purification and analysis of the enzyme from strains that cleave malodour precursor compounds:

Corynebacterium striatum Ax20 is selected to isolate and purify the enzyme responsible for the cleavage of the precursor Nα-3-hydroxy-3-methyl-hexanoyl-L-glutamine. The strain is grown during 48 h in Mueller-Hinton broth amended with 0.01% Tween 80. A total volume of 2 L of culture is harvested by centrifugation. The pellet is washed in Buffer A (50mM NaCl; 50 mM NaH2PO4/K2HPO4 buffer at pH 7) and this buffer is used throughout the whole purification procedure. The cells are disrupted mechanically by vortexing them with glass beads (425-600 µm, Sigma, St-Louis, USA) during 30 min at maximal speed. The crude cell lysate is then fractionated by precipitation with an increasing concentration of (NH₄)₂SO₄. The precipitate obtained between 50% and 80% saturation of (NH₄)₂SO₄ contained the active enzyme. This enriched sample is dissolved in Buffer A and then sequentially passed over four chromatography columns: DEAE Sepharose CL-6B anion exchange resin (Pharmacia, Uppsala, Sweden; elution with a linear gradient from 0 to 800 mM KCl); Phenyl-Sepharose hydrophobic interaction resin (Pharmacia; elution with a linear gradient from 1000 mM to 0 mM of (NH₄)₂SO₄; Mono Q strong anion exchange column on the FPLC system (Pharmacia; elution with a gradient from 0 to 800 mM KCl) and finally Mono P weak anion exchange column on the FPLC (elution with a gradient from 0-800mM KCl in a 50 mM Bis-Tris buffer instead of Buffer A). After each column separation the active fractions (determined by fluorescent activity assay with Nox-lauroyl-L-glutamine as substrate) are pooled and then desalted and concentrated by ultrafiltration (Amicon membrane YM10, Millipore, Bedford, US). The resulting active fractions after the last column separation contained



24

one major protein band with an apparent molecular weight of about 48kDa as determined by SDS-PAGE. Its effective molecular mass is determined by nano-ESI MS analysis and found to be 43365±5 Da. This enzyme retains all its activity if incubated with PMSF (Phenylmethylsulfonylfluoride, Roche Biochemicals, Mannheim, Germany) and Pefabloc SC (4-(2-aminoethyl)-benzenesulfonylfluoride, Roche Biochemicals), which are typical inhibitors for serin- and cystein proteases. On the other hand it is completely inhibited by 1 mM of EDTA and o-phenantrolin. This inhibition can be reversed by the addition of 1 mM ZnCl₂.

Example 4

Measuring inhibitory activity

Extracts of Ax 20 are prepared by mechanical disruption as described in Application Example 3.

The extract (50 μl ml corresponding to 0.2 ml initial cell culture) is added to 50 μl of Buffer A. Various concentrations of the compounds of the present invention are added in a volume of 40 μl, and after 10 min preincubation at 37°C, the reaction is and amended with 10 μl of substrate (Nα-lauroyl-L-glutamine, final concentration 50 μM). The samples are incubated for 15 min and then the reaction is stopped by adding 75 μl of Fluorescamine (2.5 ml dissolved in Acetonitrile). The resulting fluorescence resulting from derivatization of the released glutamine with fluorescamine is determined at an excitation wavelength of 381 nm and an emission wavelength of 470 nm. By comparing the samples containing compounds of the present invention with control samples with enzyme and substrate only, the inhibition (%) is calculated. Alternatively, the same assay is made either with purified enzyme from the wild-type strain (see example 3) or with a recombinantly formed enzyme produced with a strain containing an expression vector comprising a nucleic acid sequence encoding for the enzyme. Compounds of the present invention tested with this method show inhibitor activity in the nano-molar to tens of nano-molar range.

In order to evaluate enzyme activity in intact cells, stationary phase living cells of Ax20 are harvested and resuspended in Buffer A to an optical density at 600 nm of 0.25.

Inhibitory compounds are added at various concentrations, and after a preincubation of 15 min, the substrate (Nα-lauroyl-L-glutamine, 1 mM final concentration) is added. The samples are incubated for 1 h and then extracted with MTBE and HCl and analysed for released lauric acid using capillary GC. By comparing the samples containing compounds of the present invention with control samples with bacteria and substrate only, the inhibition (%) is calculated. By comparing the inhibitory capacity of the compounds on the isolated enzyme with the values obtained using intact cells, the relative uptake of the compounds by the cells can be assessed. The compounds of the present invention can cross the bacterial cell wall and cytoplasmatic membrane, and thus can have inhibitory activity in living cells at a concentration below 1 μM.

25

26

Sequence listing part of description

SEQUENCE LISTING

<110> Givaudan SA

<120> Organic compounds

<130> 30076 GB

<160> 2

<170> PatentIn version 3.1

<210> 1

<211> 399

<212> PRT

<213> Corynebacterium striatum

<400> 1

Ala Gln Glu Asn Leu Gln Lys Ile Val Asp Ser Leu Glu Ser Ser Arg

Ala Glu Arg Glu Glu Leu Tyr Lys Trp Phe His Gln His Pro Glu Met 20 25

Ser Met Gln Glu His Glu Thr Ser Lys Arg Ile Ala Glu Glu Leu Glu

tys Leu Gly Leu Glu Pro Gln Asn Ile Gly Val Thr Gly Gln Val Ala 55 60

Val The Lys Asn Gly Glu Gly Pro Ser Val Ala Phe Arg Ala Asp Phe

27

Asp Ala Leu Pro Ile Thr Glu Asn Thr Gly Leu Asp Tyr Ser Ala Asp 85 90 95

Pro Glu Leu Gly Met Met His Ala Cys Gly His Asp Leu His Thr Thr
100 105 110

Ala Leu Leu Gly Ala Val Arg Ala Leu Val Gli Asn Lys Asp Leu Trp 115 120 125

Ser Gly Thr Phe Ile Ala Val His Gln Pro Gly Glu Glu Gly Gly Gly 130 135 140

Gly Ala Arg His Met Val Asp Asp Gly Leu Ala Glu Lys Ile Ala Ala 145 150 155

Pro Asp Val Cys Phe Ala Gln His Val Phe Asn Glu Asp Pro Ala Phe 165 170 175

Gly Tyr Val Phe Thr Pro Gly Arg Phe Leu Thr Ala Ala Ser Asn Trp 180 185 190

Arg Ile His Ile His Gly Glu Gly His Gly Ser Arg Pro His Leu 195 200 205

Thr Lys Asp Pro Ile Val Val Ala Ala Ser Ile Ile Thr Lys Leu Gln 210 215 220

Thr Ile Val Ser Arg Glu Val Asp Pro Asn Glu Val Ala Val Val Thr 225 230 235 240

Val Gly Ser Ile Glu Gly Gly Lys Ser Thr Asn Ser Ile Pro Tyr Thr 245 250 255

Val Thr Leu Gly Val Asn Thr Arg Ala Ser Asn Asp Glu Leu Ser Glu 260 265 270

Tyr Val Gln Asn Ala Ile Lys Arg Ile Val Ile Ala Glu Cys Gln Ala 275 280 285 ·

Ala Gly Ile Glu Glu Glu Pro Glu Phe Glu Tyr Leu Asp Ser Val Pro 290 295 300

Ala Val Ile Asn Asp Glu Asp Leu Thr Glu Gln Leu Met Ala Gln Phe

28

305

('

310

315

320

Arg Glu Phe Phe Gly Glu Asp Gln Ala Val Glu Ile Pro Pro Leu Ser 325 330 335

Gly Ser Glu Asp Tyr Pro Phe Ile Pro Asn Ala Trp Gly Val Pro Ser

Val Met Trp Gly Trp Ser Gly Phe Ala Ala Gly Ser Amp Ala Pro Gly 355 360 365

Asn His Thr Asp Lys Phe Ala Pro Glu Leu Pro Asp Ala Leu Glu Arg 370 375 380

Gly Thr Gln Ala Ile Leu Val Ala Ala Pro Trp Leu Met Lys 385 390 395

<210> 2

<211> 1212

<212> DNA

<213> Corynebacterium striatum

<400> 2 aatogggtca tggcacagga aaatttgcaa aagattgtag atagtetoga gteeteeege 60 goggaacgcg aagaactgta caagtggtte caccagoacc cggaaatgtc gatgcaggag cacgaaacct ccaagcgcat cgcagaagag ctagagaagc tcggccttga gccgcagaac 180 ateggegtga cegggeaggt egeggtaate aagaaeggtg aaggeeegag egtggeattt egtgeggaet ttgatgoett geegateace gagaacaceg ggetggatta eteggoggat 300 cccgagctgg gcatgatgca cgcctgcggc cacgatttgc acaccactgc cctactcggc 360 goggtgcgcg cgctggtgga gaacaaggac ctgtggtccg gcaccttcat cgcagtccac 420 caacccggrg aggaaggceg cggcggggcc cgccacatgg tggacgaogg cctcgcggag 480 aagategegg egeeggatgt gtgtttegee eageacgtgt teaacgaaga eceegeettt ggctacgtgt teacccccgg ccggtttcta acggcggcgt cgaactggag aatccacatc 600 cacggogagg gcggacacgg ttcccgtccg cacctgacca aggacccgat tgtggtggcg geetegatea traccaaget geagacgatt gretecegeg aagregatee gaargaggre

geagtggtea cegteggete eategaggge ggeaagteea ceaactegat ceegtabace 780 gteaccteg gegtgaacac cegageeter aacgatgage teteogagta egtecagaac 840 gocatoaago goatogtoat ogoggagtgo caggotgoag goatogaaca ggagooggaa 900 ttegagtace tggacteagt ecoggeegtg ateaacgaeg aggateteac egaacagete 960 atggcgcagt tocgggagtt cttcggcgag gaccaggcgg tagagattcc gcccotgtcc 1020 ggcagcgagg actacccct catteegaac gcctggggcg tgccgagtgt gatgtgggga 1080 tggtccggct tcgccgcagg ttctgaogca ccgggcaatc acaccgacaa gttcgccccc 1140 gagettecag atgoectega acgeggeace caggecatte tggtggccgc cgcgccotgg 1200 1212 ttgatgaagt ga

·30076GB

30

Claims

1. A compound of formula (I)

wherein

```
is a substituted alkyl, henzyl or allyl residue selected from the group consisting of
R
nonyl;
```

4,4,4-trifluoro-propyl;

2-methyl-4-phenyl-butyl;

4-trifluoromethyl-phenyl;

pentafluorophenyl;

4-fluoro-phenyl;

naphthalene-2-yl;

biphenyl-2-yl:

4-tert-butyl-phenyl;

5,5,7,8,8-pentamethyl-5,6,7,8-tetrahydro-naphthalene-2-yl;

5,5,8,8-tetramethyl-5,6,7,8-tetrahydro-naphthalen-2-yl;

1,1,3,3-tetramethyl-indan-5-yl;

styryl;

2,6-dimethyl-hept-1,5-dienyl;

2-(4-tert-Butyl-phenyl)-1-methyl-vinyl;

2-(4-lsopropyl-phenyl)-1-methyl-vinyl;

1-Methyl-3-(2,2,3-trimethyl-cyclopent-3-enyl)-propenyl;

2-(4-Isobutyl-phenyl)-1-methyl-vinyl;

2-(2-isopropyl-phenyl)-1-methyl-ethenyl;

2-phenyl-ethyl;

cyclohexyl-methyl;

41 1 824 29 26;



30076GB

31

- 2,2-dimethyl-propyl;
- 2-pentafluorophenyl-ethyl;
- 3-phenyl-propyl; heptyl; and
- 4-isopropyl-cyclohex-1-enyl.
- 2. A composition comprising a body odour-suppressing quantity of a compound claimed in claim 1.
- Composition according to claim 2 wherein the compound is present in amounts of about 0.01 to 0.5 % by weight.
- 4. Composition according to claim 2 or claim 3 selected from cosmetic and personal care products, in particular deo-sticks, roll-ons, pump-sprays, aerosols, deodorant soaps, powders, solutions, gels, creams, sticks, balms and lotions.
- 5. Use of a compound or composition to inhibit an enzyme in its ability to cleave compounds contained in sweat into short-chained, branched fatty acids, which enzyme is produced in bacteria of the genus *Corynebacteria*, which bacteria has been deposited at the International Depository Authority DSMZ- Deutsche Sammlung von Mikrooganismen und Zellkulturen GmbH, D-38124 Braunschweig under the Accession Number DSM 14267.
- 6. Use of a compound or composition to inhibit an enzyme in its ability to cleave compounds contained in sweat into short-chained, branched fatty acids, which enzyme is defined by an amino acid sequence set forth as SEQ ID No. 1.
- 7. A method of suppressing axillary malodour comprising the step of providing a composition for application to a person in need of treatment, said composition containing an inhibitor compound and a dermatologically acceptable vehicle therefor, said compound being selected from a screening of compounds for activity in the inhibition of the enzyme.

32

Abstract

::

Inhibitors of axillary malodour having the formula

This Page is Inserted by IFW Indexing and Scanning Operations and is not part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images include but are not limited to the items checked:

BLACK BORDERS

IMAGE CUT OFF AT TOP, BOTTOM OR SIDES

FADED TEXT OR DRAWING

BLURRED OR ILLEGIBLE TEXT OR DRAWING

SKEWED/SLANTED IMAGES

COLOR OR BLACK AND WHITE PHOTOGRAPHS

REFERENCE(S) OR	EXHIBIT(S) SUBMITTED A	RE POOR QUALITY	
□ OTHER:			

IMAGES ARE BEST AVAILABLE COPY.

☐ LINES OR MARKS ON ORIGINAL DOCUMENT

☐ GRAY SCALE DOCUMENTS

As rescanning these documents will not correct the image problems checked, please do not report these problems to the IFW Image Problem Mailbox.